## SEQUENCE LISTING

## <110> VLAAMS INTERUNIVERSITAIR INSTITUUT VOOR BIOTECHNOL <120> Nucleic Acid Binding of Multi-Zinc Finger Transcription Factors <130> 2676-5174US <140> US/10/028,396 <141> 2001-12-21 <150> 99202068.5 <151> 1999-06-25 <150> PCT/EP00/05582 <151> 2000-06-09 <160> 49 <170> PatentIn version 3.1 <210> 1 <211> 5 <212> DNA <213> Artificial <220> <221> misc feature <223> Description of Artificial Sequence: Portion of bait for screening <400> 1 5 cacct <210> 2 <211> 6 <212> DNA <213> Artificial <220> <221> misc feature <223> Description of Artificial Sequence: portion of bait for screening

<400> 2

cacctg

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<210> 3
<211> 5
<212> DNA
<213> Artificial
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<400> 3
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aggtg
<210> 4
<211> 7
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<223> Description of Artificial Sequence: consensus element for binding
      of MyT1, NZF-1 and NZF-3
<400> 4
                                          7
aaagttt
<210> 5
<211> 52
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<220>
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<223> Description of Artificial Sequence: complex consensus sequence
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<222> (16)..(43)
<223> nucleotides 16-43 represent a spacer sequence wherein any one, more,
       or all of nucleotides 16-43 my be present or absent
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<400> 5

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<210> 6
<211> 30
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<220>
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<400> 6
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ccacctgaaa gaatccctga gaattcacag
<210> 7
<211> 30
<212> DNA
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<223> Description of Artificial Sequence: primer SIP1 CZF2Mut
<400> 7
                                                  30
gggtcctaca gttcatctat cagcagcaag
<210> 8
<211> 30
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<213> Artificial
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<223> Description of Artificial Sequence: primer SIP1 NZF4Mut
<400> 8
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caccacctta tcgagtcctc gaggctgcac
<210> 9
<211> 30
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<212> DNA <213> Artificial

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<210> 10
<211> 50
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<223> Description of Artificial Sequence: probe Xbra-WT
<400> 10
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atccaggcca cctaaaatat agaatgataa agtgaccagg tgtcagttct
<210> 11
<211> 50
<212> DNA
<213> Artificial
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<223> Description of Artificial Sequence: probe Xbra-D
<400> 11
atccaggcca cctaaaatat agaatgataa agtgaccaga tgtcagttct
                                                          50
<210> 12
<211> 23
<212> DNA
<213> Artificial
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<223> Description of Artificial Sequence: probe Xbra-E
<400> 12
                                                23
taaagtgacc aggtgtcagt tct
<210> 13
<211> 27
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<220>
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<223> Description of Artificial Sequence: probe Xbra-F
<400> 13
                                                  27
atccaggcca cctaaaatat agaatga
<210> 14
<211> 50
<212> DNA
<213> Artificial
<220>
<221> misc feature
<223> Description of Artificial Sequence: probe Rdm + Xbra-E
<400> 14
caatttagag tactgtgtac ttgggagtaa agtgaccagg tgtcagttct
                                                         50
<210> 15
<211> 53
<212> DNA
<213> Artificial
<220>
<221> misc feature
<223> Description of Artificial Sequence: probe Xbra-F + AREB6
<400> 15
                                                            53
atccaggcca cctaaaatat agaatgaggc tcagacaggt gtagaattcg gcg
<210> 16
<211> 53
<212> DNA
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<400> 16
caatttagag tactgtgtac-ttgggagggc-tcagacaggt-gtagaattcg-gcg
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<211> 50
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<223> Description of Artificial Sequence: probe Xbra-J
<400> 17
gcacaggcca cctaaaatat agaatgataa agtgaccagg tgtcagttct
                                                           50
<210> 18
<211> 50
<212> DNA
<213> Artificial
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<223> Description of Artificial Sequence: probe Xbra-K
<400> 18
atcactgcca cctaaaatat agaatgataa agtgaccagg tgtcagttct
                                                          50
<210> 19
<211> 50
<212> DNA
<213> Artificial
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<223> Description of Artificial Sequence: probe Xbra-L
<400> 19
atccagtaaa cctaaaatat agaatgataa agtgaccagg tgtcagttct
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<210> 20
<211> 50
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<212> DNA

<213>-Artificial

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<210> 21
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<400> 21
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<210> 22
<211> 50
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<400> 22
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atccaggcca cctaaccgat agaatgataa agtgaccagg tgtcagttct
<210> 23
<211> 50
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<400> 23
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<210> 24
<211> 50
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<223> Description of Artificial Sequence: probe Xbra-Q
<400> 24
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atccaggcca cctaaaatat atcctgataa agtgaccagg tgtcagttct
<210> 25
<211> 50
<212> DNA
<213> Artificial
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<400> 25
                                                           50
atccaggcca cctaaaatat agaagtctaa agtgaccagg tgtcagttct
<210> 26
<211> 50
<212> DNA
<213> Artificial
<220>
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<400> 26
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<210> 27
<211> 50
<212> DNA
<213> Artificial
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<223> Description of Artificial Sequence: probe Xbra-Z
<400> 27
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<210> 28
<211> 47
<212> DNA
<213> Artificial
<220>
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<223> Description of Artificial Sequence: probe Xbra-B
<400> 28
atccaggcca cctatataga atgataaagt gaccaggtgt cagttct
                                                        47
<210> 29
<211> 47
<212> DNA
<213> Artificial
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<223> Description of Artificial Sequence: probe Xbra-C
<400> 29
                                                        47
atccaggcca cctaaaatat agaatgatgt gaccaggtgt cagttct
<210> 30
<211> 40
<212> DNA
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<223> Description of Artificial Sequence: probe Xbra-U
<400> 30
atccaggcca cctaaaatat agtgaccagg tgtcagttct
                                                      40
<210> 31
<211> 46
<212> DNA
<213> Artificial
<220>
<221> misc feature
<223> Description of Artificial Sequence: probe Xbra-EE
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<210> 33 <211> 50 <212> DNA

<213> Artificial

<220>

<221> misc\_feature

<223> Description of Artificial Sequence: probe Xbra-FrF

<400> 33

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<210> 34 <211> 50

<212> DNA <213> Artificial

<220>

<221> misc\_feature

<223> Description of Artificial Sequence: probe Xbra-V

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<210> 35

<211> 50

<212> DNA

<213> Artificial

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<400> 35
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<210> 36
<211> 60
<212> DNA
<213> Artificial
<220>
<221> misc feature
<223> Description of Artificial Sequence: probe alfa-4I-WT (alfa-4-integrin)
<400> 36
gcagggcaca cetggattge attagaatga gacteactae ceagtteagg tgtgttgegt
<210> 37
<211> 60
<212> DNA
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<223> Description of Artificial Sequence: probe alfa-4I-A (alfa-4-integrin)
<400> 37
                                                                60
geagggeaca cetggattge attagaatga gacteactae ceagtteaga tgtgttgegt
<210> 38
<211> 60
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<223> Description of Artificial Sequence: probe alfa-4I-B (alfa-4-integrin)
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<210> 39 <211> 70

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<400> 39
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                                           70
ctggctgcag
<210> 40
<211> 70
<212> DNA
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<221> misc feature
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                                           70
ctggctgcag
<210> 41
<211> 70
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<213> Artificial
<220>
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<400> 41
tggccggcag atgaaccete agccaatcag cggtacgggg ggcggtgete cgggggeteae
                                                                 60
                                           70
ctggctgcag
<210> 42
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<213> Artificial

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|---|--|
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| _ | sequence (-341/+41)  |
|   | <400> 42   |
|   | acaaaagaac tcagccaagt g 21   |
|   | ucunuuguuo tougoouugt g  |
|   | <210> 43   |
|   | <211> 18   |
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|   | <213> Artificial   |
|   | <220>  |
|   | <221> misc feature   |
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|   | sequence (-341/+41)  |
|   |  |
|   | <400> 43   |
|   | cegeaagete acaggtge 18   |
|   | 2010s 44   |
|   | <210> 44<br><211> 26   |
|   | <211> 20<br><212> DNA  |
|   | <213> Artificial   |
|   | ~215~ Attitiviai   |
|   | <220>  |
|   | <221> misc feature   |
|   | <223> Description of Artificial Sequence: forward primer E-box1                    |
|   |  |
|   |  |
|   | <400> 44 getgtggcg geagatgaac ceteag 26  |
|   | getgtggccg gcagatgaac cetcag 26  |
|   | <210> 45   |
|   | <211> 26   |
|   | <212> DNA  |
|   | <213> Artificial   |
|   |  |
|   | <220>  |
|   | <221> misc_feature <223> Description of Artificial Sequence: reverse primer E-box1 |
|   | ~223/ Description of Authorit Sequence. Teverse printer 2 cont                     |
|   | <400> 45   |
|   | ctgagggttc atctgccggc cacagc 26  |
|   |  |

| <211> 24   |                       |  |
|--|-----------------------|--|
| ~211° 2¬   |                       |  |
| <212> DNA  |                       |  |
| <213> Artificial   |                       |  |
|  |                       |  |
| <220>  |                       |  |
| <221> misc_feature   |                       |  |
| <223> Description of Artificial Sequence:  | forward primer E-box3 |  |
| <100> 16   |                       |  |
| <400> 46   | 24                    |  |
| geteeggget eatetggetg eage 24  |                       |  |
| <210> 47   |                       |  |
| <211> 25   |                       |  |
| <212> DNA  |                       |  |
| <213> Artificial   |                       |  |
|  |                       |  |
| <220>  |                       |  |
| <221> misc_feature   |                       |  |
| <223> Description of Artificial Sequence:  | reverse primer E-box3 |  |
|  |                       |  |
| <400> 47   |                       |  |
| gctgcagcca gatgagcccc ggagc  | 25                    |  |
| 212  |                       |  |
| <210> 48   |                       |  |
| <211> 27   |                       |  |
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| <212> DNA  |                       |  |
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<220>
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<222> (26)
<223> n is a spacer and may be any nucleotide
<400> 49
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28